therapeutically active agent may be associated with a container holding a flowable carrier component (e.g. a container may hold fibrinogen and C3).

SEQUENCE LISTING

~~(1) GENERAL INFORMATION:~~

(i) APPLICANT: LISA MCKERRACHER

(ii) TITLE OF INVENTION: Methods for making and delivering Rho-antangonist tissue adhesive formulations to the injured mammalian central and peripheral nervous systems and uses thereof

(iii) NUMBER OF SEQUENCES: 3

~~(iv) CORRESPONDENCE ADDRESS:~~

~~(A) ADRESSEE: BROULLETTE KOSIE~~

~~(B) STREET: 1100 RENE LESVEQUE BLVD WEST~~

~~(C) PROV/STATE: QUEBEC~~

~~(D) COUNTRY: CANADA~~

~~(E) POSTAL/ZIP CODE: H3B 5C9~~

~~(v) COMPUTER READABLE FORM:~~

~~(A) MEDIUM TYPE: Floppy disk~~

~~(B) COMPUTER: IBM PC compatible~~

~~(C) OPERATING SYSTEM: PC-DOS/MS-DOS~~

(D) SOFTWARE: ~~ASCII (TEXT)~~ PatentIn version 3.1

~~(vi) CURRENT APPLICATION DATA:~~

~~(A) APPLICATION NUMBER:~~

~~(B) FILING DATE:~~

~~(C) CLASSIFICATION:~~

~~(vii) ATTORNEY/AGENT INFORMATION:~~

~~(A) NAME: RONALD S. KOSIE~~

~~(B) REGISTRATION NO.: 28,814~~

(C) REFERENCE/DOCKET NO.: ~~06447-003-US-2~~ 06447-011

~~(D) TEL. NO.: (514) 397-8500~~

~~(E) FAX NO.: (514) 397-8515~~

(2) INFORMATION FOR SEQ ID NO: 1:

~~(i) SEQUENCE CHARACTERISTICS:~~

(A) LENGTH:

(B) TYPE:

~~(C) STRANDEDNESS:~~

~~(D) TOPOLOGY:~~

~~(ii) MOLECULE TYPE:~~

~~(v) FRAGMENT TYPE:~~

~~(vii) ORIGINAL SOURCE~~:

(A) ORGANISM:

~~(vii) IMMEDIATE SOURCE:~~

~~(ix) FEATURE:~~

(A) NAME/KEY:

(B) LOCATION:

(D) OTHER INFORMATION:

(x) PUBLICATION INFORMATION:

(A) AUTHORS:

(B) TITLE:

(C) JOURNAL:

(D) VOLUME:

(E) ISSUE:

(F) PAGES:

(G) DATE:

(H) DOCUMENT NO.:

(I) FILING DATE:

(J) PUBLICATION DATE:

(K) RELEVANT RESIDUES IN SEQ ID NO:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

GTG GCG ACC CTT CCC AAA TCG GAT CTG GTT CCG CGT GGA TCC TCT AGA

5 10 15

GTC GAC CTG CAG GCA TGC AAT GCT TAT TCC ATT AAT CAA AAG GCT TAT

20 25 30

TCA AAT ACT TAC CAG GAG TTT ACT AAT ATT GAT CAA GCA AAA GCT TGG

35 40 45

GGT AAT GCT CAG TAT AAA AAG TAT GGA CTA AGC AAA TCA GAA AAA GAA

50 55 60

GCT ATA GTA TCA TAT ACT AAA AGC GCT AGT GAA ATA AAT GGA AAG CTA

65 70 75 80

AGA CAA AAT AAG GGA GTT ATC AAT GGA TTT CCT TCA AAT TTA ATA AAA

85 90 95

CAA GTT GAA CTT TTA GAT AAA TCT TTT AAT AAA ATG AAG ACC CCT GAA

100 105 110

AAT ATT ATG TTA TTT AGA GGC GAC GAC CCT GCT TAT TTA GGA ACA GAA

115 120 125

TTT CAA AAC ACT CTT CTT AAT TCA AAT GGT ACA ATT AAT AAA ACG GCT

130 135 140

TTT GAA AAG GCT AAA GCT AAG TTT TTA AAT AAA GAT AGA CTT GAA TAT

145 150 155 160

GGA TAT ATT AGT ACT TCA TTA ATC AAT GTT TCT CAA TTT GCA GGA AGA

165 170 175

CCA ATT ATT ACA AAA TTT AAA GTA GCA AAA GGC TCA AAG GCA GGA TAT

180 185 190

ATT GAC CCT ATT AGT GCT TTT CAG GGA CAA CTT GAA ATG TTG CTT CCT

195 200 205

AGA CAT AGT ACT TAT CAT ATA GAC GAT ATG AGA TTG TCT TCT GAT GGT

210 215 220

AAA CAA ATA ATA ATT ACA GCA ACA ATG ATG GGC ACA GCT ATC AAT CCT

225 230 235 240

AAA TAA

(2) INFORMATION FOR SEQ ID NO: 2:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH:

(B) TYPE:

(C) STRANDEDNESS:

(D) TOPOLOGY:

(vi) ORIGINAL SOURCE:

(A) ORGANISM:

~~(ix) FEATURE:~~

~~(D) OTHER INFORMATION:~~

~~(xi) SEQUENCE DESCRIPTION~~: SEQ ID NO: 2:

GGATCCTCTA GAGTCGACCT GCAGGCATGC AATGCTTATT CCATTAATCA ~~50~~
AAAGGCTTAT TCAAATACTT ACCAGGAGTT TACTAATATT GATCAAGCAA ~~100~~
AAGCTTGGGG TAATGCTCAG TATAAAAAGT ATGGACTAAG CAAATCAGAA ~~150~~
AAAGAAGCTA TAGTATCATA TACTAAAAGC GCTAGTGAAA TAAATGGAAA ~~200~~
GCTAAGACAA AATAAGGGAG TTATCAATGG ATTTCCTTCA AATTTAATAA ~~250~~
AACAAGTTGA ACTTTTAGAT AAATCTTTTA ATAAAATGAA GACCCCTGAA ~~300~~
AATATTATGT TATTTAGAGG CGACGACCCT GCTTATTTAG GAACAGAATT ~~350~~
TCAAAACACT CTTCTTAATT CAAATGGTAC AATTAATAAA ACGGCTTTTG ~~400~~
AAAAGGCTAA AGCTAAGTTT TTAAATAAAG ATAGACTTGA ATATGGATAT ~~450~~
ATTAGTACTT CATTAATGAA TGTTTCTCAA TTTGCAGGAA GACCAATTAT ~~500~~
TACAAAATTT AAAGTAGCAA AAGGCTCAAA GGCAGGATAT ATTGACCCTA ~~550~~
TTAGTGCTTT TCAGGGACAA CTTGAAATGT TGCTTCCTAG ACATAGTACT ~~600~~
TATCATATAG ACGATATGAG ATTGTCTTCT GATGGTAAAC AAATAATAAT ~~650~~
TACAGCAACA ATGATGGGCA CAGCTATCAA TCCTAAATAA

(2) INFORMATION FOR SEQ ID NO: 3:

~~(i) SEQUENCE CHARACTERISTICS~~:

(A) LENGTH:

(B) TYPE:

~~50~~

~~(C) STRANDEDNESS~~:

~~(D) TOPOLOGY:~~

~~(vi) ORIGINAL SOURCE~~:

(A) ORGANISM:

~~(ix) FEATURE:~~

~~(D) OTHER INFORMATION~~:

~~(xi) SEQUENCE DESCRIPTION~~: SEQ ID NO: 3:

GSSRVDLQAC NAYSINQKAY SNTYQEFTNI DQAKAWGNAQ YKKYGLSKSE ~~50~~
KEAIVSYTKS ASEINGKLRQ NKGVINGFPS NLIKQVELLD KSFNKMKTPE ~~100~~
NIMLFXGDDP AYLGTEFQNT LLNSNGTINK TAFEKAKAKF LNXDRLEYGY ~~150~~
ISTSLMNVSQ FAGRPIITKF KVAKGSKAGY IDPISAFQGQ LEMLLPRHST ~~200~~
YHIDDMRLSS DGKQIIITAT MMGTAINPK